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<p>(21) International Application Number: PCT/US99/20207</p> <p>(22) International Filing Date: 2 September 1999 (02.09.99)</p> <p>(30) Priority Data: 60/099,068 3 September 1998 (03.09.98) US 60/135,835 24 May 1999 (24.05.99) US</p> <p>(71) Applicant (<i>for all designated States except US</i>): LOMA LINDA UNIVERSITY [US/US]; Loma Linda, CA 92350 (US).</p> <p>(72) Inventors; and (75) Inventors/Applicants (<i>for US only</i>): SZALAY, Aladar, A. [US/US]; 7327 Fairwood Lane, Highland, CA 92346 (US). WANG, Yubao [CN/US]; 24931 Prospect Avenue, Loma Linda, CA 92354 (US). WANG-PRUSKI, Gefu [CA/CA]; 114 Hillcrest Avenue, Truro, Nova Scotia B2N 4L3 (CA).</p> <p>(74) Agents: SHELDON, Jeffrey, G. et al.; Sheldon &amp; Mak, 9th floor, 225 South Lake Avenue, Pasadena, CA 91101 (US).</p>	<p>(81) Designated States: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).</p> <p>Published <i>With international search report.</i></p>	
<p>(54) Title: METHOD FOR STUDYING PROTEIN INTERACTIONS <i>IN VIVO</i></p> <p>(57) Abstract</p> <p>A method for determining whether a first protein interacts with a second protein within a living cell. The method comprises providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore within the cell. The complexed first protein and the complexed second protein are allowed to come into proximity to each other within the cell. Then, any fluorescence from the acceptor fluorophore resulting from luminescence resonance energy transfer from the donor luciferase is detected, where fluorescence from the acceptor fluorophore indicates that the first protein has interacted with the second protein.</p>		

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**METHOD FOR STUDYING PROTEIN INTERACTIONS *IN VIVO***

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**BACKGROUND**

The study of interactions between proteins in living cells is often necessary to understand the proteins' functions and their mechanisms of action. These interactions are currently studied using immuno-precipitation, the yeast two hybrid method, and  $\beta$ -gal complementation method.

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However, these methods are associated with several disadvantages. For example, these methods are associated with false positives. Second, they do not permit the determination of quantitative information regarding the interactions. Further, they do not allow for *in vivo* real time monitoring of the interactions.

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Therefore, it would be advantageous to have another method of studying interactions between proteins *in vivo*, which does not have these disadvantages. Further preferably, the method could be used with a wide variety of proteins and in a wide variety of living cells. Also preferably, the method could be used to determine the interactions between molecules other than proteins.

**SUMMARY**

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According to one embodiment of the present invention, there is provided a

method for determining whether a first protein interacts with a second protein within a living cell. The method comprises providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore within the cell. The donor luciferase is capable of luminescence resonance energy transfer to the acceptor fluorophore when the first protein is in proximity to the second protein. Then, the complexed first protein and the complexed second protein are allowed to come into proximity to each other within the cell. Next, any fluorescence from the acceptor fluorophore is detected. Fluorescence of the acceptor fluorophore resulting from luminescence resonance energy transfer from the donor luciferase to acceptor fluorophore the indicates that the first protein has interacted with the second protein.

In a preferred embodiment, providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore comprises genetically engineering DNA and transferring the genetically engineered DNA to the living cell causing the cell to produce the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore. In a particularly preferred embodiment, the cell which is provided with the first protein complexed to a donor luciferase and the cell which is provided with the second protein complexed to an acceptor fluorophore are mammalian cells.

In another preferred embodiment, the donor luciferase provided is *Renilla* luciferase. In yet another preferred embodiment, the acceptor fluorophore provided is an *Aequorea* green fluorescent protein.

In a particularly preferred embodiment, the detection of acceptor fluorophore fluorescence is performed using spectrofluorometry.

#### DESCRIPTION

The present invention includes a method for determining whether a first protein interacts with a second protein in a living cell using luminescent resonance energy transfer (LRET). Luminescence resonance energy transfer results from the transfer of excited state energy from a donor luciferase to an acceptor fluorophore. In order for LRET to occur, there must be an overlap between the emission spectrum of the donor luciferase and the excitation spectrum of the acceptor fluorophore.

The efficiency of luminescence resonance energy transfer is dependent on the distance separating the donor luciferase and the acceptor fluorophore, among other variables.

Generally, significant energy transfers occur only where the donor luciferase and acceptor fluorophore are less than about 80 Å of each other. This short distance is considerably less than the distance needed between for optical resolution between two entities using conventional microscopy. Therefore, detecting luminescence resonance energy transfer between a donor luciferase and an acceptor fluorophore indicates that the donor luciferase and acceptor fluorophore have come within the distance needed for LRET to occur, that is less than about 80 Å of each other.

The present invention utilizes luminescence resonance energy transfer to determine whether an interaction takes place between a first protein and a second protein in a living cell. This is accomplished by complexing a first protein to the donor luciferase and complexing the second protein to the acceptor fluorophore and placing the complexed first protein and the complexed second protein in the cell under conditions suitable for an interaction between the first protein and the second protein to take place. If the first protein interacts with the second protein, the donor luciferase will come close enough to the acceptor fluorophore for luminescence resonance energy transfer to take place and the acceptor fluorophore will fluoresce. Detection of fluorescence from the acceptor fluorophore will, thereby, indicate that the first protein has interacted with the second protein.

Advantageously, this method allows for the detection of interaction between the first protein and the second protein even though the interaction cannot be detected by optical methods such as conventional microscopy.

There are several advantages of using luminescent resonance energy transfer to detect the interaction between two proteins according to the present invention. First, the specific labeling of the proteins in living cells can be achieved through genetic engineering methods where the introduction of fluorescent dyes into living cells is very difficult. Further, fluorescent dyes photobleach quickly while light emission of a luciferase such as *Renilla* luciferase originates from an enzymatic reaction that is relatively stable if substrate and oxygen are supplemented.

As used in this disclosure, "complexing a first protein to the donor luciferase" refers to joining the donor luciferase to the first protein in a manner that the donor luciferase and the first protein stay in essentially the same proximity to one another during interaction between the first protein and the second protein. Similarly, "complexing a second protein to

the acceptor fluorophore" refers to joining the acceptor fluorophore to the second protein in a manner that the acceptor fluorophore and the second protein stay in essentially the same proximity to one another during interaction between the first protein and the second protein. Such complexing can be done, for example, by genetically engineering the cell to produce a fusion protein containing the donor luciferase and first protein, and the acceptor fluorophore and the second protein.

In a preferred embodiment, the present invention uses *Renilla* luciferase as the donor luciferase and "humanized" *Aequorea* green fluorescent protein ('humanized' GFP) as the acceptor fluorophore. *Renilla* luciferase is a 34 kDa enzyme purified from *Renilla reniformis*. The enzyme catalyzes the oxidative decarboxylation of coelenterazine in the presence of oxygen to produce blue light with an emission wavelength maximum of 471 nm. *Renilla* luciferase was used as the donor luciferase because it requires an exogenous substrate rather than exogenous light for excitation. This, advantageously, eliminates background noise from an exogenous light source and from autofluorescence, and allows easy and accurate quantitative determination of light production.

'Humanized' GFP is a 27 kDa protein fluorophore that has an excitation maximum at 480 nm. It has a single amino acid difference from wild-type *Aequorea* green fluorescent protein. 'Humanized' GFP was chosen as the acceptor fluorophore because its excitation spectrum overlaps with the emission spectra of *Renilla* luciferase. Additionally, emissions from 'humanized' GFP can be visualized in living cells. Further, 'humanized' GFP is expressed well in the mammalian cells transfected with 'humanized' GFP cDNA that were used to demonstrate this method.

The method for determining whether a first protein interacts with a second protein according to the present invention was demonstrated as follows. In summary, insulin-like growth factor binding protein 6 (IGFBP 6) and insulin-like growth factor II (IGF-II) were selected as the first protein and second protein. IGFBP 6 is a protein known to have a marked binding affinity for IGF-II.

The *Renilla* luciferase cDNA was fused to IGFBP 6 cDNA and 'humanized' GFP cDNA was fused to IGF-II cDNA. Living cells were transfected with the fused cDNAs and the fusion proteins were expressed. Cell extracts were produced and mixed. The substrate for the *Renilla* luciferase moiety of the fused *Renilla* luciferase-IGFBP 6 protein

was added. Finally, fluorescence from the 'humanized' GFP moiety of the fused 'humanized' GFP-IGF-II protein was detected. Demonstration one method according to the present invention will now be described in greater detail.

**A) The Cloning of Fused IGFBP-6 cDNA to *Renilla* Luciferase cDNA; Fused IGF-II cDNA to 'humanized' GFP cDNA; and Fused Insulin cDNA to 'humanized' GFP cDNA:**

First, three fused cDNAs were produced: 1) fused IGFBP-6 cDNA and *Renilla* luciferase cDNA; 2) fused IGF-II cDNA and 'humanized' GFP cDNA; and 3) fused insulin cDNA and 'humanized' GFP cDNA. IGFBP-6 cDNA, SEQ ID NO:1, GenBank accession number M69054, encoded IGFBP-6, SEQ ID NO:2, which was used as the first protein.

*Renilla* luciferase cDNA, SEQ ID NO:3, GenBank accession number M63501, encoded *Renilla* luciferase, SEQ ID NO:4, which was used as the donor luciferase. IGF-II cDNA, SEQ ID NO:5, encoded IGF-II, SEQ ID NO:6, which was used as the second protein.

'Humanized' GFP cDNA, SEQ ID NO:7, GenBank accession number U50963, encoded 'humanized' GFP, SEQ ID NO:8, which was used as the acceptor fluorophore. Insulin cDNA, SEQ ID NO:9, accession number AH002844, encoded insulin, SEQ ID NO:10.

Insulin, fused to 'humanized' GFP, was used as a control protein because insulin is homologous to IGF-II, but it does not bind to IGFBP-6. The IGFBP-6 cDNA, SEQ ID NO:1, IGF-II cDNA, SEQ ID NO:5, and insulin cDNA, SEQ ID NO:9, were modified using PCR as follows.

First, the cDNA of prepro-IGF-II carried on an EcoRI fragment was cloned into pBluescript KS (+) II vector. The insert was sequenced using T7 and T3 primers and confirmed to contain the known cDNA sequence of prepro-IGF-II. The 5' end of the IGF-II precursor was connected to the T7 promoter in the pBluescript KS (+) II vector. An IGF-II 3' primer was designed to generate a Notice of Allowance restriction site, to remove the D and E domains of prepro-IGF-II, and to maintain the Notice of Allowance fragment of the 'humanized' GFP in frame with the open reading frame of IGF-II.

Next, the IGF-II fragment was amplified with PCR using the T7 promoter primer and the IGF-II 3' primer. The PCR-amplified IGF-II fragment was digested by EcoRI and Not I and cloned into pCDNA3.1 (+) vector (Invitrogen, Carlsbad, CA, US) producing pCDNA-IGF-II. Then, the Notice of Allowance fragment of the 'humanized' GFP was inserted into the Not I site of pCDNA-IGF-II producing pC-IGF-II-GFP.

The cDNA for precursor of insulin, which contained a signal peptide the B, C and A domains, was modified in a manner corresponding to the IGF-II fragment, above. The 'humanized' GFP cDNA was then linked to the 3' end of the modified insulin cDNA to produce pC-INS-GFP.

5                   Finally, IGFBP 6 cDNA was amplified by PCR from a plasmid named Rat-tagged human IGFBP6. The stop codon of IGFBP 6 was removed and the open reading frame of IGFBP 6 was in frame with *Renilla* luciferase cDNA from pCEP4-RUC (Mayerhofer R, Langridge WHR, Cormier MG and Szalay AA. *Expression of recombinant Renilla luciferase in transgenic plants results in high levels of light emission*. The Plant  
10   Journal 1995;7;1031-8). The linking of the *Renilla* luciferase cDNA to the 3' end of modified IGFBP 6 cDNA produced pC-IGFBP 6-RUC.

The sequences of the insert DNA fragments from all the constructs were verified by DNA sequencing analysis. Qiagen Maxi Plasmid Kit (Qiagen, Inc., Valencia, CA) was used for the purification of plasmid DNA.

15   **B) Transient Transfection of Mammalian Cells With pC-IGF-II-GFP, pC-INS-GFP and pC-IGFBP 6-RUC Using the Calcium Phosphate Precipitation Method:**

Next, mammalian cells were transfected with the cloned fusion DNAs. First, COS-7 cells (African green monkey kidney cell, American Type Culture Collection CRL 1651) were grown at 37 C in Dulbecco's Modified Eagle Medium (DMEM) with  
20   L-Glutamine supplemented with 10% fetal bovine serum and antibiotic antimycotic solution containing a final concentration of penicillin 100 unit/ml, streptomycin 100 mg/ml and amphotericin B 250 ng/ml (Sigma-Aldrich Co., St. Louis, MO, US) in 5% CO<sub>2</sub>. Groups of 1x10<sup>6</sup> of these cells were plated the day before transfection and were approximately 50% to 60% confluent at the time of transfection.

25                   Forty mg of each plasmid fusion DNA were precipitated and resuspended into Dulbecco's Phosphate Buffered Saline Solution and the plasmid fusion DNAs was introduced into mammalian cells using the standard calcium phosphate precipitation method. Transfection efficiency was estimated by fluorescence microscopy after 24 hours. The number of green fluorescent cells per plate were comparable in plates of pC-IGF-II-GFP  
30   DNA transfected cells, pC-INS-GFP DNA transfected cells and cells transfected with a plasmid DNA containing GFP only, which was used as a positive control.

**C) Confirmation of Expression of Fusion Proteins:**

Twenty-four hours after DNA transfection using DNA calcium phosphate precipitation method, individual plasmid DNA transfected COS-7 cells were visualized using fluorescence microscopy by detection of GFP fluorescence. pC-IGF-II-GFP and pC-INS-GFP transfected cells showed similar fluorescence patterns typical of secretory protein translocated through ER to Golgi. The pC-IGFBP 6-RUC transfected cells did not fluoresce. However, the pC-IGFBP 6-RUC transfected cells did show luminescence using a low light imaging system after the addition of coelenterazine.

Further, the presence of fusion proteins IGF-II-GFP and IGFBP 6-RUC, having the expected molecular weights of about 36 kDa and 56 kDa, respectively, were detected using immunoblot analysis. This confirmed the presence of both fusion proteins in the transiently transfected cells.

**D) Detection of Protein Interactions by Spectrofluorometry:**

Having confirmed the presence of the expected fusion proteins IGF-II-GFP and IGFBP 6-RUC, and the function of the donor luciferase and acceptor fluorophore, cell extracts from these transiently transfected cells were used to carry out a protein binding assay based on energy transfer between the *Renilla* luciferase and 'humanized' GFP moieties of the fusion proteins. Forty-eight hours after calcium transfection, the COS cells were washed twice with PBS and harvested using a cell scraper in luciferase assay buffer containing 0.5 M NaCl, 1 mM EDTA and 0.1 M potassium phosphate at a pH 7.5. The harvested cells were sonicated 3 times for 10 seconds with an interval of 10 seconds using a Fisher Model 550 Sonic Dismembrator (Fisher Scientific, Pittsburgh, PA, US) to produce cell extracts.

Next, the cell extracts containing IGF-II-GFP and IGFBP 6-RUC were mixed and 0.1  $\mu$ g of coelenterazine was immediately added. Spectrofluorometry was performed using a SPEX FluoroMax® (Instruments S.A., Inc., Edison, NJ). The spectrum showed a single emission peak at 471 nm, which corresponds to the known emission of *Renilla* luciferase.

Following the first spectrofluorometry, the mixtures were kept at room temperature for 30 minutes and the spectra were traced again after fresh coelenterazine was added. The trace at 30 minutes showed two peaks with emission maximum at 471 nm and 503 nm. The spectrofluorometry of the cell extracts was carried out at a longer time, but the

spectral pattern did not change over time.

Control cell extract mixtures from cells transfected with pC-INS-GFP and pC-IGFBP 6-RUC were made similarly and their spectra traced. The traces showed only one peak at 471 nm, which corresponds to the emission peak of *Renilla* luciferase. The spectral pattern did not change over time.

Therefore, these data demonstrated that IGFBP 6 and IGF-II interacted but that insulin and IGFBP 6 did not interact.

In addition to the above disclosed examples, protein-protein interactions were also detected by the detection of LRET using corresponding methods in *E. coli* cells and mammalian cells which were co-transformed.

Although the present invention has been discussed in considerable detail with reference to certain preferred embodiments, other embodiments are possible. For example, the interaction between molecules other than proteins could be studied by corresponding methods. Such other molecules could be provided to the living cell by diffusion, infusion, and incorporation or by other means. Further, fusion proteins produced from genetically engineered living cells could have post translational changes, such as the addition of sugar moieties, before their interactions are studied. Also, living cells can be visualized using these methods by spectrofluorometry by low light image analysis in cells, colonies and tissues. Additionally, high through put screening of colonies can be accomplished using the present methods combined with cell sorting and low light video analysis of micro titre dishes or multiple array detection. Therefore, the spirit and scope of the appended claims should not be limited to the description of preferred embodiments contained herein.

**WHAT IS CLAIMED IS:**

1. A method for determining whether a first protein interacts with a second protein within a living cell, the method comprising:

a) providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore within the cell;

b) placing the complexed first protein and the complexed second protein in proximity to each other within the cell; and

c) detecting any fluorescence from the acceptor fluorophore;

where the donor luciferase is capable of luminescence resonance energy transfer to the acceptor fluorophore when the first protein is in proximity to the second protein; and

where fluorescence of the acceptor fluorophore resulting from luminescence resonance energy transfer from the donor luciferase indicates that the first protein has interacted with the second protein.

2. The method of claim 1, where providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore comprises genetically engineering DNA and transferring the genetically engineered DNA to the living cell causing the cell to produce the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore.

3. The method of claim 1, where the cell provided with the first protein complexed to a donor luciferase is a mammalian cell.

4. The method of claim 1, where the cell provided with the second protein complexed to a acceptor fluorophore is a mammalian cell.

5. The method of claim 1, where the donor luciferase provided is *Renilla* luciferase.

6. The method of claim 1, where the acceptor fluorophore provided is a green fluorescent protein.

7. The method of claim 1, where the acceptor fluorophore provided is an *Aequorea* green fluorescent protein.

8. The method of claim 1, where detecting any fluorescence from the donor luciferase is performed using spectrofluorometry.

9. A method for determining whether a first molecule interacts with a second molecule within a living cell, the method comprising:

a) providing the first molecule complexed to a donor luciferase and the second molecule complexed to an acceptor fluorophore within the cell;

b) placing the complexed first molecule and the complexed second molecule in proximity to each other within the cell; and

5 c) detecting any fluorescence from the acceptor fluorophore;

where the donor luciferase is capable of luminescence resonance energy transfer to the acceptor fluorophore when the first molecule is in proximity to the second molecule; and

where fluorescence of the acceptor fluorophore resulting from luminescence resonance energy transfer from the donor luciferase indicates that the first molecule has interacted with  
10 the second molecule.

10. The method of claim 9, where the first molecule is a first protein and where the second molecule is a second protein; and

where providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore comprises genetically engineering DNA and  
15 transferring the genetically engineered DNA to the living cell causing the cell to produce the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore.

11. The method of claim 10, where the cell provided with the first protein complexed to a donor luciferase is a mammalian cell.

20 12. The method of claim 10, where the cell provided with the second protein complexed to a acceptor fluorophore is a mammalian cell.

13. The method of claim 9, where the donor luciferase provided is *Renilla* luciferase.

14. The method of claim 9, where the acceptor fluorophore provided is a green fluorescent protein.

25 15. The method of claim 9, where the acceptor fluorophore provided is a *Aequorea* green fluorescent protein.

16. The method of claim 9, where detecting any fluorescence from the donor luciferase is performed using spectrofluorometry.

17. A method for determining whether a first protein interacts with a second protein,  
30 the method comprising:

a) providing the first protein complexed to a donor luciferase and the second protein

complexed to an acceptor fluorophore;

b) placing the complexed first protein and the complexed second protein in proximity to each other; and

c) detecting any fluorescence from the acceptor fluorophore;

5 where the donor luciferase is capable of luminescence resonance energy transfer to the acceptor fluorophore when the first protein is in proximity to the second protein; and

where fluorescence of the acceptor fluorophore resulting from luminescence resonance energy transfer from the donor luciferase indicates that the first protein has interacted with the second protein.

10 18. The method of claim 17, where providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore comprises genetically engineering DNA and transferring the genetically engineered DNA to a living cell causing the cell to produce the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore.

15 19. The method of claim 18, where the cell provided with the first protein complexed to a donor luciferase is a mammalian cell.

20. The method of claim 18, where the cell provided with the second protein complexed to a acceptor fluorophore is a mammalian cell.

20 21. The method of claim 17, where the donor luciferase provided is *Renilla* luciferase.

22. The method of claim 17, where the acceptor fluorophore provided is a green fluorescent protein.

23. The method of claim 17, where the acceptor fluorophore provided is an *Aequorea* green fluorescent protein.

25 24. The method of claim 17, where detecting any fluorescence from the donor luciferase is performed using spectrofluorometry.

25. A method for determining whether a first molecule interacts with a second molecule, the method comprising:

30 a) providing the first molecule complexed to a donor luciferase and the second molecule complexed to an acceptor fluorophore;

b) placing the complexed first molecule and the complexed second molecule in

proximity to each other; and

c) detecting any fluorescence from the acceptor fluorophore;

where the donor luciferase is capable of luminescence resonance energy transfer to the acceptor fluorophore when the first molecule is in proximity to the second molecule; and

5        where fluorescence of the acceptor fluorophore resulting from luminescence resonance energy transfer from the donor luciferase indicates that the first molecule has interacted with the second molecule.

26. The method of claim 25, where the first molecule is a first protein and where the second molecule is a second protein; and

10        where providing the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore comprises genetically engineering DNA and transferring the genetically engineered DNA to a living cell causing the cell to produce the first protein complexed to a donor luciferase and the second protein complexed to an acceptor fluorophore.

15        27. The method of claim 26, where the cell provided with the first protein complexed to a donor luciferase is a mammalian cell.

28. The method of claim 26, where the cell provided with the second protein complexed to a acceptor fluorophore is a mammalian cell.

20        29. The method of claim 25, where the donor luciferase provided is *Renilla* luciferase.

30. The method of claim 25, where the acceptor fluorophore provided is a green fluorescent protein.

31. The method of claim 25, where the acceptor fluorophore provided is a *Aequorea* green fluorescent protein.

25        32. The method of claim 25, where detecting any fluorescence from the donor luciferase is performed using spectrofluorometry.

## SEQUENCE LISTING

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Wang, Yubao

Wang-Pruski, Gefu

Loma Linda University

&lt;120&gt; Method for Studying Protein Interactions in Vivo

&lt;130&gt; 11785-3PCT

&lt;140&gt; filed herewith

&lt;141&gt; 1999-09-02

&lt;150&gt; 60/135,835

&lt;151&gt; 1999-05-24

&lt;150&gt; 60/099,068

&lt;151&gt; 1998-09-03

&lt;160&gt; 10

&lt;170&gt; PatentIn Ver. 2.0

&lt;210&gt; 1

&lt;211&gt; 918

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (3)..(719)

&lt;400&gt; 1

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Leu Leu Ala Ala Ser Pro Gly Gly Ala Leu Ala Arg Cys Pro Gly Cys

20 25 30

ggg caa ggg gtg cag gcg ggt tgt cca ggg ggc tgc gtg gag gag gag 143

Gly Gln Gly Val Gln Ala Gly Cys Pro Gly Gly Cys Val Glu Glu Glu

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Ser Pro Asp Gly Asn Gly Ser Ser Ser Cys Pro Thr Gly Ser Ser Gly			
225	230	235	
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caacaaaaaa ccgaggccct caatccacct tcagggcccg ccccatgggc ccctcaccgc 839  
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<212> PRT

<213> Homo sapiens

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Gly Gly Ser Pro Ala Glu Gly Cys Ala Glu Ala Glu Gly Cys Leu Arg  
 50 55 60

Arg Glu Gly Gln Glu Cys Gly Val Tyr Thr Pro Asn Cys Ala Pro Gly  
 65 70 75 80

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 85 90 95

Leu Leu Gly Arg Gly Arg Cys Leu Pro Ala Arg Ala Pro Ala Val Ala  
 100 105 110

Glu Glu Asn Pro Lys Glu Ser Lys Pro Gln Ala Gly Thr Ala Arg Pro  
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Gln Asp Val Asn Arg Arg Asp Gln Gln Arg Asn Pro Gly Thr Ser Thr  
 130 135 140

Thr Pro Ser Gln Pro Asn Ser Ala Gly Val Gln Asp Thr Glu Met Gly  
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Pro Cys Arg Arg His Leu Asp Ser Val Leu Gln Gln Leu Gln Thr Glu  
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Val Tyr Arg Gly Ala Gln Thr Leu Tyr Val Pro Asn Cys Asp His Arg  
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Gly Phe Tyr Arg Lys Arg Gln Cys Arg Ser Ser Gln Gly Gln Arg Arg  
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 Ile Thr Gly Pro Gln Trp Ala Arg Cys Lys Gln Met Asn Val Leu  
 15 20 25 30

gat tca ttt att aat tat tat gat tca gaa aaa cat gca gaa aat gct 147  
 Asp Ser Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn Ala  
 35 40 45

gtt att ttt tta cat ggt aac gcg gcc tct tct tat tta tgg cga cat 195  
 Val Ile Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp Arg His  
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gtt gtg cca cat att gag cca gta gcg cgg tgt att ata cca gat ctt 243  
 Val Val Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile Pro Asp Leu  
 65 70 75

att ggt atg ggc aaa tca ggc aaa tct ggt aat ggt tct tat agg tta 291  
 Ile Gly Met Gly Lys Ser Gly Lys Ser Gly Asn Gly Ser Tyr Arg Leu  
 80 85 90

ctt gat cat tac aaa tat ctt act gca tgg ttt gaa ctt ctt aat tta 339  
 Leu Asp His Tyr Lys Tyr Leu Thr Ala Trp Phe Glu Leu Leu Asn Leu  
 95 100 105 110

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Phe His Tyr Ser Tyr Glu His Gln Asp Lys Ile Lys Ala Ile Val His
      130              135              140

gct gaa agt gta gta gat gtg att gaa tca tgg gat gaa tgg cct gat 483
Ala Glu Ser Val Val Asp Val Ile Glu Ser Trp Asp Glu Trp Pro Asp
      145              150              155

att gaa gaa gat att gcg ttg atc aaa tct gaa gaa gga gaa aaa atg 531
Ile Glu Glu Asp Ile Ala Leu Ile Lys Ser Glu Glu Gly Glu Lys Met
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atg aga aag tta gaa cca gaa gaa ttt gca gca tat ctt gaa cca ttc 627
Met Arg Lys Leu Glu Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe
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aaa gag aaa ggt gaa gtt cgt cgt cca aca tta tca tgg cct cgt gaa 675
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atc ccg tta gta aaa ggt ggt aaa cct gac gtt gta caa att gtt agg 723
Ile Pro Leu Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg
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att gaa tcg gat cca gga ttc ttt tcc aat gct att gtt gaa ggc gcc 819
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aag aag ttt cct aat act gaa ttt gtc aaa gta aaa ggt ctt cat ttt 867
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tcg caa gaa gat gca cct gat gaa atg gga aaa tat atc aaa tcg ttc 915
Ser Gln Glu Asp Ala Pro Asp Glu Met Gly Lys Tyr Ile Lys Ser Phe
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 35 40 45  
 Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp Arg His Val Val  
 50 55 60  
 Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile Pro Asp Leu Ile Gly  
 65 70 75 80  
 Met Gly Lys Ser Gly Lys Ser Gly Asn Gly Ser Tyr Arg Leu Leu Asp  
 85 90 95  
 His Tyr Lys Tyr Leu Thr Ala Trp Phe Glu Leu Leu Asn Leu Pro Lys  
 100 105 110  
 Lys Ile Ile Phe Val Gly His Asp Trp Gly Ala Cys Leu Ala Phe His  
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 Tyr Ser Tyr Glu His Gln Asp Lys Ile Lys Ala Ile Val His Ala Glu  
 130 135 140  
 Ser Val Val Asp Val Ile Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu

145                                      150                                      155                                      160  
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 Glu Asn Asn Phe Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg  
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 Lys Leu Glu Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu  
    195                                      200                                      205  
 Lys Gly Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro  
    210                                      215                                      220  
 Leu Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn Tyr  
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    245                                      250                                      255  
 Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly Ala Lys Lys  
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 Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu His Phe Ser Gln  
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Cys Gly Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys Gly Asp Arg  
 35 40 45

Gly Phe Tyr Phe Ser Arg Pro Ala Ser Arg Val Ser Arg Arg Ser Arg  
 50 55 60

Gly Ile Val Glu Glu Cys Cys Phe Arg Ser Cys Asp Leu Ala Leu Leu  
 65 70 75 80

Glu Thr Tyr Cys Ala Thr Pro Ala Lys Ser Glu Arg Asp Val Ser Thr  
 85 90 95

Pro Pro Thr Val Leu Pro Asp Asn Phe Pro Arg Tyr Pro Val Gly Lys  
 100 105 110

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Gly Leu Pro Ala Leu Leu Arg Ala Arg Arg Gly His Val Leu Ala Lys  
 130 135 140

Glu Leu Glu Ala Phe Arg Glu Ala Lys Arg His Arg Pro Leu Ile Ala  
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Leu Pro Thr Gln Asp Pro Ala His Gly Gly Ala Pro Pro Glu Met Ala  
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Ser Asn Arg Lys  
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&lt;211&gt; 717

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

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&lt;222&gt; (1)..(717)

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ggg gaa ggt gat gcc aca tac gga aag ctc acc ctg aaa ttc atc tgc  144
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acc atc ttt ttc aaa gat gac ggg aac tac aag acc cgc gct gaa gtc  336
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aaa gat ccc aac gaa aag aga gac cac atg gtc ctg ctg gag ttt gtg 672  
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Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe  
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Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln  
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His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg  
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Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val  
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Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile  
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Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn  
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Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly  
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Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val  
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Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro  
 180 185 190

Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser  
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tgg gga cct gac cca gcc gca gcc ttt gtg aac caa cac ctg tgc ggc 96  
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cag gtg gag ctg ggc ggg ggc cct ggt gca ggc agc ctg cag ccc ttg 240  
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<213> Homo sapiens

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Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe  
 35 40 45

Phe Tyr Thr Pro Lys Thr Arg Arg Glu Ala Glu Asp Leu Gln Val Gly  
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Gln Val Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu  
 65 70 75 80

Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys  
 85 90 95

Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn  
 100 105 110

## INTERNATIONAL SEARCH REPORT

International Application No.

PCT/US99/20207

**A. CLASSIFICATION OF SUBJECT MATTER**

IPC(6) : Please See Extra Sheet.

US CL : Please See Extra Sheet.

According to International Patent Classification (IPC) or to both national classification and IPC

**B. FIELDS SEARCHED**

Minimum documentation searched (classification system followed by classification symbols)

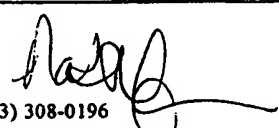
U.S. : Please See Extra Sheet.

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched  
NONEElectronic data base consulted during the international search (name of data base and, where practicable, search terms used)  
WEST, CAS ONLINE**C. DOCUMENTS CONSIDERED TO BE RELEVANT**

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y, E	US 5,976,796 A (SZALAY et al.) 02 November 1999, col. 1, line 60-col. 3, line 17.	1-32
Y	US 5,491,084 A (CHALFIE et al.) 13 February 1996, col. 1, lines 36-61.	1-32
Y	US 5,418,155 A (CORMIER et al.) 23 May 1995, col. 2, line 59-col. 10, line 9.	1-32
Y, P	US 5,891,646 A (BARAK et al.) 06 April 1999, col. 2, line 6-col. 4, line 47.	1-32
A	US 5,866,348 A (SCHEIRER) 02 February 1999, col. 2, line 15-col. 3, line 27.	1-32

☒ Further documents are listed in the continuation of Box C.
 ☐ See patent family annex.

* Special categories of cited documents:	*T* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
*A* document defining the general state of the art which is not considered to be of particular relevance	*X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
*B* earlier document published on or after the international filing date	*Y* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
*L* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	*A* document member of the same patent family
*O* document referring to an oral disclosure, use, exhibition or other means	
*P* document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search 06 NOVEMBER 1999	Date of mailing of the international search report <b>19 NOV 1999</b>
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Facsimile No. (703) 305-3230	Authorized officer PENSEE T. DO  Telephone No. (703) 308-0196

## INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US99/20207

## C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	US 4,604,364 A (KOSAK) 05 August 1986, col. 3, line 26-col. 8, line 14.	1-32
A	US 4,318,707 A (LITMAN et al.) 09 March 1982, col. 3, line 1-col. 7, line 14.	1-32
A	US 5,683,888 A (CAMPBELL) 04 November 1997, col. 2, line 10-col. 4, line 50.	1-32
A, P	US 5,811,238 A (STEMMER et al.) 22 September 1998, col. 4, line 11-col. 7, line 35.	1-32

# INTERNATIONAL SEARCH REPORT

International Application No.

PCT/US99/20207

## A. CLASSIFICATION OF SUBJECT MATTER:

IPC (6):

C12Q 1/66, 1/00; C12P 21/06; C07H 21/04; G01N 33/53, 33/567, 33/573, 33/58, 33/60, 33/531, 33/533, 21/00, 21/76, 21/75, 31/00

## A. CLASSIFICATION OF SUBJECT MATTER:

US CL :

435/4, 7.1, 7.2, 7.21, 7.4, 7.91, 8, 68.1, 69.1, 70.1, 70.3, 127, 231, 232, 326, 331, 366, 375, 968, 972; 436/543, 546, 164, 172, 800, 805, 905; 536/23.5; 424/9.6

## B. FIELDS SEARCHED

Minimum documentation searched

Classification System: U.S.

435/4, 7.1, 7.2, 7.21, 7.4, 7.91, 8, 68.1, 69.1, 70.1, 70.3, 127, 231, 232, 326, 331, 366, 375, 968, 972; 436/543, 546, 164, 172, 800, 805, 905; 536/23.5; 424/9.6